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RAW SEQUENCE LISTING  
PATENT APPLICATION: US/10/044,643

DATE: 01/30/2002  
TIME: 15:14:41

Input Set : A:\Cura-241.app  
Output Set: N:\CRF3\01302002\J044643.raw

Q.S

3 <110> APPLICANT: Majumder, Kumud  
 4 Vernet, Corine  
 5 Casman, Stacie J  
 6 Wolenc, Adam R  
 7 Spaderna, Steven K  
 8 Padigaru, Muralidhara  
 9 Mishnu, Vishun S  
 10 Tchernev, Velizar T  
 11 Spytek, Kimberly A  
 12 Li, Li  
 13 Baumgartner, Jason C  
 14 Gusev, Vladimir  
 16 <120> TITLE OF INVENTION: Novel Proteins and Nucleic Acids Encoding Same  
 18 <130> FILE REFERENCE: 15966-748  
 C--> 20 <140> CURRENT APPLICATION NUMBER: US/10/044,643  
 C--> 21 <141> CURRENT FILING DATE: 2002-01-11  
 23 <150> PRIOR APPLICATION NUMBER: 60/193,664  
 24 <151> PRIOR FILING DATE: 2000-03-31  
 26 <150> PRIOR APPLICATION NUMBER: 60/194,614  
 27 <151> PRIOR FILING DATE: 2000-04-05  
 29 <150> PRIOR APPLICATION NUMBER: 60/195,063  
 30 <151> PRIOR FILING DATE: 2000-04-06  
 32 <150> PRIOR APPLICATION NUMBER: 60/195,066  
 33 <151> PRIOR FILING DATE: 2000-04-06  
 35 <150> PRIOR APPLICATION NUMBER: 60/195,067  
 36 <151> PRIOR FILING DATE: 2000-04-06  
 38 <150> PRIOR APPLICATION NUMBER: 60/195,068  
 39 <151> PRIOR FILING DATE: 2000-04-06  
 41 <150> PRIOR APPLICATION NUMBER: 60/195,069  
 42 <151> PRIOR FILING DATE: 2000-04-06  
 44 <150> PRIOR APPLICATION NUMBER: 60/195,070  
 45 <151> PRIOR FILING DATE: 2000-04-06  
 47 <150> PRIOR APPLICATION NUMBER: 60/195,510  
 48 <151> PRIOR FILING DATE: 2000-04-06  
 50 <150> PRIOR APPLICATION NUMBER: 60/219,855  
 51 <151> PRIOR FILING DATE: 2000-07-21  
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 54 <151> PRIOR FILING DATE: 2000-07-27  
 56 <150> PRIOR APPLICATION NUMBER: 60/221,325  
 57 <151> PRIOR FILING DATE: 2000-07-28  
 59 <150> PRIOR APPLICATION NUMBER: 60/224,588  
 60 <151> PRIOR FILING DATE: 2000-08-11  
 62 <150> PRIOR APPLICATION NUMBER: 60/239,613

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66 <151> PRIOR FILING DATE: 2001-01-18
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75 <151> PRIOR FILING DATE: 2001-01-30
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79 <170> SOFTWARE: PatentIn Ver. 2.1
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83 <212> TYPE: DNA
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89 gtttctgtt ccacatggaa accttggaaac ccagcactgt ttacccatcc aattttggcgc 180
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96 tcatgttccca ggtttggatc ttatgtcccc tcggccatcat ctatttttgc tccttcgaaga 600
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119 Asn Gly Val Ala Leu Cys Gly Phe Cys Phe His Met Lys Thr Trp Lys
120 35 40 45
122 Pro Ser Thr Val Tyr Leu Phe Asn Leu Ala Val Ala Asp Phe Leu Leu
123 50 55 60
125 Met Ile Cys Leu Pro Phe Arg Thr Asp Tyr Tyr Leu Arg Arg Arg His
126 65 70 75 80

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Input Set : A:\Cura-241.app  
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128 Trp Ala Phe Gly Asp Ile Pro Cys Arg Val Gly Leu Phe Thr Leu Ala  
 129 85 90 95  
 131 Met Asn Arg Ala Gly Ser Ile Val Phe Leu Thr Val Val Ala Ala Asp  
 132 100 105 110  
 134 Arg Tyr Phe Lys Val Val His Pro His His Ala Val Asn Thr Ile Ser  
 135 115 120 125  
 137 Thr Arg Val Ala Ala Gly Ile Val Cys Thr Leu Trp Ala Leu Val Ile  
 138 130 135 140  
 140 Leu Gly Thr Val Tyr Leu Leu Leu Glu Asn His Leu Cys Val Gln Gln  
 141 145 150 155 160  
 143 Thr Ala Val Ser Cys Glu Ser Phe Ile Met Glu Ser Ala Asn Gly Trp  
 144 165 170 175  
 146 His Asp Ile Met Phe Gln Leu Glu Phe Met Pro Leu Gly Ile Ile  
 147 180 185 190  
 149 Leu Phe Cys Ser Phe Lys Ile Val Trp Ser Leu Arg Arg Arg Gln Gln  
 150 195 200 205  
 152 Leu Ala Arg Gln Ala Arg Met Lys Lys Ala Thr Arg Phe Ile Met Val  
 153 210 215 220  
 155 Val Ala Ile Val Phe Ile Thr Cys Tyr Leu Pro Ser Val Ser Ala Arg  
 156 225 230 235 240  
 158 Leu Tyr Phe Leu Trp Thr Val Pro Ser Ser Ala Cys Asp Pro Ser Val  
 159 245 250 255  
 161 His Gly Ala Leu His Ile Thr Leu Ser Phe Thr Tyr Met Asn Ser Met  
 162 260 265 270  
 164 Leu Asp Pro Leu Val Tyr Tyr Phe Ser Ser Pro Ser Phe Pro Lys Phe  
 165 275 280 285  
 167 Tyr Asn Lys Leu Lys Ile Cys Ser Leu Lys Pro Lys Gln Pro Gly His  
 168 290 295 300  
 170 Ser Lys Thr Gln Arg Pro Glu Met Pro Ile Ser Asn Leu Gly Arg  
 171 305 310 315 320  
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 174 325 330 335  
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 188 gtttctgtt ccacatgaag acctggaaagc ccagcaactgt ttaccttttc aatttgcgc 180  
 189 tgccgtgattt ctccttcttgc atctgcgttc cttttccggac agacttattac ctcagacgtta 240  
 190 gacactgggc tttttggggac atttccttgc gagggtggct cttcacggtt gccatgaaaca 300  
 191 gggccgggag catcggtttc cttacgggtgg tggctggcga caggtatttc aaagtggttc 360  
 192 accggccacca cgcgcgttgc acatcttccca cccgggtggc ggctggcata gctctgacccc 420  
 193 tggggccctt ggttcatcttgc ggaaacgtgt atctttgttgc ggagaaccat ctctgggtgc 480  
 194 aagagacggc cgttcttctgtt yagagtttca tcatgggtc ggccaaatggc tggcatgaca 540  
 195 tcatgttccca gctggaggttc ttatgtcccc tcggcatcat cttatgttgc tccttcaaga 600

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Input Set : A:\Cura-241.app

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196 ttgtttggag cctgaggcgg aggacagcgc tggccagaca ggctcgatg aagaaggcga 660  
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 198 cttagactcta ttccctctgg acggtgccct cgagtgctg cgatccctct gcccatgggg 780  
 199 ccctgcacat aaccctcagc ttccacctaca ttaacagcat gctggatccc ctggtgatt 840  
 200 attttcaag cccctctttt cccaaattct acaacaagct caaaatctgc agtctgaaac 900  
 201 ccaaggcggc aggacactca aaaaacacada ggcggaaaga gatgccaatt tggaaacctcg 960  
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 231 <213> ORGANISM: Homo sapiens  
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 235 Val Met Pro Pro Leu Leu Ile Val Ala Phe Val Leu Gly Ala Leu Asp  
 236 20 25 30  
 237 Asn Gly Val Ala Leu Cys Gly Phe Cys Phe His Met Lys Thr Trp Lys  
 238 35 40 45  
 239 Pro Ser Thr Val Tyr Leu Phe Asn Leu Ala Val Ala Asp Phe Leu Leu  
 240 50 55 60  
 241 Met Ile Cys Leu Pro Phe Arg Thr Asp Tyr Tyr Leu Arg Arg Arg His  
 242 65 70 75 80  
 243 Trp Ala Phe Gly Asp Ile Pro Cys Arg Val Gly Leu Phe Thr Leu Ala  
 244 85 90 95  
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Input Set : A:\Cura-241.app  
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260 Arg Tyr Phe Lys Val Val His Pro His His Ala Val Asn Thr Ile Ser  
261 115 120 125  
263 Thr Arg Val Ala Ala Gly Ile Val Cys Thr Leu Trp Ala Leu Val Ile  
264 130 135 140  
266 Leu Gly Thr Val Tyr Leu Leu Leu Glu Asn His Leu Cys Val Gln Glu  
267 145 150 155 160  
269 Thr Ala Val Ser Cys Glu Ser Phe Ile Met Glu Ser Ala Asn Gly Trp  
270 165 170 175  
272 His Asp Ile Met Phe Gln Leu Glu Phe Phe Met Pro Leu Gly Ile Ile  
273 180 185 190  
275 Leu Phe Cys Ser Phe Lys Ile Val Trp Ser Leu Arg Arg Arg Gln Gln  
276 195 200 205  
278 Leu Ala Arg Gln Ala Arg Met Lys Lys Ala Thr Arg Phe Ile Met Val  
279 210 215 220  
281 Val Ala Ile Val Phe Ile Thr Cys Tyr Leu Pro Ser Val Ser Ala Arg  
282 225 230 235 240  
284 Leu Tyr Phe Leu Trp Thr Val Pro Ser Ser Ala Cys Asp Pro Ser Val  
285 245 250 255  
287 His Gly Ala Leu His Ile Thr Leu Ser Phe Thr Tyr Met Asn Ser Met  
288 260 265 270  
290 Leu Asp Pro Leu Val Tyr Tyr Phe Ser Ser Pro Ser Phe Pro Lys Phe  
291 275 280 285  
293 Tyr Asn Lys Leu Lys Ile Cys Ser Leu Lys Pro Lys Gln Pro Gly His  
294 290 295 300  
296 Ser Lys Thr Gln Arg Pro Glu Glu Met Pro Ile Ser Asn Leu Gly Arg  
297 305 310 315 320  
299 Arg Ser Cys Ile Ser Val Ala Asn Ser Phe Gln Ser Gln Ser Asp Gly  
300 325 330 335  
302 Gln Trp Asp Pro His Ile Val Glu Trp His  
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314 ctgcgcgttc cggactgtgt gtggggactgt gggctggagt tgcggacggc cgcgcgcgcca 180  
315 ggcatccccc cggggcggcgg cggggcggagc agcgcggacaca cagaggccccg ggtgcggatt 240  
316 ctcatcagcg tggtgtactg tggtgtgc ggccctgggt tggcgccaa cttgtctgg 300  
317 ctctacccgtc tgaagacatc gcggcgctgg cgcacgttctt ctatcaacctt cttcgtcacc 360  
318 aaccttggcgc tgacgcgactt tcagtttgcg ctccacccgtc cttcttgggc ggtggagaaac 420  
319 gctcttgact tcaaatggcc ctccggcaag gccatgtgttca agatctgttc catgtgtacg 480  
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322 ggccggggcc tggggggacac ctgtgttgc tggcccaagg cgcgtgtgtt gttggatctgg 660  
323 gctttggccgc cgcgtggccctc gtcgtccatgttccatccaccggcgttgcgtgttgc 720

Use of n or Xaa has been detected in the Sequence Listing.  
The following sequence listing to insure a corresponding  
explanation is presented in the <210> to <223> fields of  
each sequence using n or Xaa.

VERIFICATION SUMMARY

PATENT APPLICATION: US/10/044,643

DATE: 01/30/2002

TIME: 15:14:42

Input Set : A:\Cura-241.app

Output Set: N:\CRF3\01302002\J044643.raw

L:20 M:270 C: Current Application Number differs, Replaced Current Application Number

L:21 M:271 C: Current Filing Date differs, Replaced Current Filing Date

L:2854 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:53

L:3541 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:63